



# Polyamines: Metabolism, Regulation, and Functions in Crop Abiotic Stress Tolerance

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## Abstract

Polyamines are small, positively charged, organic compounds containing more than two amino groups. They are omnipresent in plants and produced during various metabolic processes. Environmental fluctuations owing to greenhouse gases, pollution, deforestation, and global warming are known to hamper plants' normal growth, production, and developmental processes causing various forms of abiotic stresses such as drought, salinity, heat, cold, osmolarity, etc. Polyamines are considered important to plants as they provide support in maintaining normal growth and development of plants even during stressed conditions. They play crucial role whether present at endogenous levels or fortified exogenously to plants utilizing plant tissue culture or genetic transformation techniques. Although they are regarded important for plants, still their mode of action and regulation during plant stress conditions is still not well understood. In this chapter, their endogenous production, mode of action and regulation is described at length so as to facilitate a broader and clearer picture to the researchers to understand the importance of them in combating various abiotic stresses in plants.

## Keywords

Polyamines · Abiotic stress tolerance · Putrescine · Spermidine · Spermine

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## 19.1 Introduction

Polyamines (PAs) are polycationic aliphatic amines. They are present in plants abundantly and take part in major plant growth and development processes. Spermidine (Spd), spermine (Spm) and putrescine (Put) are common polyamines apparently involved in plant responses to microbial symbionts that are critical for plant nutrition (El Ghachtoul et al. 1996), and are involved in molecular signaling events in interactions between plants and pathogens (Martin-Tanguy 1987). PAs are important for plant growth development and environmental stresses as well (Evans and Malmberg 1989; Galston and Kaur-Sawhney 1990). PAs are synthesized from the amino acids ornithine or arginine. Their capacities to neutralize acids and act as antioxidant agents—as well as their roles in membrane/cell wall stabilization—make them indispensable for normal functioning of cells. DNA has an overall negative charge due to phosphate group attached to it (Basu et al. 1990; Pohjanpelto and Hölttä 1996), so PAs tend to bind with DNA, thereby providing stability to the DNA helix (Beigbeder 1995; Tassoni et al. 1996) and pectic polysaccharides (D'Oraci and Bagni 1987). They have been shown to be involved in protein phosphorylation (Ye et al. 1994), post-transcriptional modifications (Mehta et al. 1994), and conformational transition of DNA (Basu et al. 1990). There is direct evidence that they are essential for growth and development in prokaryotes and eukaryotes (Tabor and Tabor 1984; Heby and Persson 1990; Tiburcio et al. 1990; Slocum 1991). PAs are considered to be signaling molecules, especially in stress situations (Evans and Malmberg 1989; Galston and Kaur-Sawhney 1995).

Naturally occurring PAs play pivotal role in plant metabolism such as effects on the tolerance mechanism to abiotic stresses (Meloni et al. 2003; Zeid 2004; Jiuju and Shirong 2005; Duan et al. 2007; Roychoudhury et al. 2011).

Abiotic stresses such as soil salinity, cold, frost injury, drought, acidity, and heavy metal affect plant growth, development, and productivity. Since these conditions produce stress and result in extensive losses to many agriculturally important crops, they have been the main subject of intense research. The accumulation of some functional substances, such as compatible solute, protective proteins, and polyamines, is an important element of the physiological and biochemical response to the stressful conditions. Polyamines have been proposed as a new category of plant growth regulators which are active in physiological processes, such as embryogenesis, cell division, morphogenesis, and development (Bais and Ravishankar 2002; Liu et al. 2006a). Though the physiological significance of polyamines in stress is not thoroughly understood, much progress has been made. The focus in the present paper is on progress concerning the involvement and potential role of polyamines in plant responses to abiotic stresses.

## 19.2 Polyamine Biosynthesis and Metabolism in Plants

Polyamine biosynthesis in plants is documented. Put is produced either directly from ornithine-by-ornithine decarboxylase (ODC, EC 4.1.1.17) or indirectly from arginine-by-arginine decarboxylase (ADC, EC 4.1.1.19) with two intermediates, agmatine and *N*-carbamoyl putrescine, and two corresponding biosynthetic enzymes, agmatine iminohydrolase (EC 3.5.3.12) and *N*-carbamoylputrescine amidohydrolase (EC 3.5.1.53) (Malmberg et al. 1998; Martin-Tanguy 2001). Put is converted into Spd via spermidine synthase (SPDS, EC 2.5.1.16) with the addition of an aminopropyl moiety provided by decarboxylated *S*-adenosylmethionine (dcSAM), which is catalyzed by *S*-adenosylmethionine decarboxylase (SAMDC, EC 4.1.1.50) using *S*-adenosylmethionine (SAM) as the substrate. Similarly, Spm is produced from Spd via spermine synthase (SPMS, EC 2.5.1.22) with the same aminopropyl moiety rendered by dcSAM. Apart from biosynthesis, polyamine degradation plays a crucial role in cellular polyamine titers regulation, which is primarily ascribed to two amine oxidases, diamine oxidase (DAO, EC 1.4.3.6), and polyamine oxidase (PAO, EC 1.5.3.11). DAO catalyzes the oxidation of Put to give pyrroline, which is further metabolized to *g*-aminobutyric acid (Cona et al. 2006) and PAO, and acts as a catalyst in the conversion of Spd and Spm to pyrroline and 1-(3 aminopropyl)-pyrroline, respectively, along with 1,3-diaminopropane in plants (Martin-Tanguy 2001; Šebela et al. 2001).

## 19.3 Polyamines Transport in Plants

It is estimated that PAs can be transported in long distances; the presence of large amounts of PAs has been observed in exudates of xylem and phloem sap; the vacuolar membrane had the highest capacity to transport them (Vladimir and Shevyakova 2007).

As per a study carried out in carrot cells, PAs (Put and Spd) flow through the cell by a transmembrane electrochemical gradient. Other research with maize roots, whose application of Put was performed exogenously, indicated that Put is transported through the plasma membrane by a process regulated by a protein carrier (Kusano et al. 2008). This carrier TPO1 of PAs is mainly located in vacuolar membrane and plasma membrane. Among the four PA carriers, those which are encoded by TPO2 and TPO3 are specific for Spm, while for Put, Spd and Spm are encoded by TPO1 and TPO4 (Uemura et al. 2005). TPO1 is dependent of pH (Uemura et al. 2005); in some research with lichen (*Evernia prunastri*), it was shown that the uptake of PAs depends on this variable (Kakkar et al. 1998). Uptake of Spr and Spd is performed at alkaline pH (pH = 8.0), while inhibition of only Spd was observed at acidic pH (pH = 5.0), next to the pH of the vacuoles found internally. This suggests that TPO1 carrier acts as a catalyst for polyamine excretion at acidic pH (Uemura et al. 2005). Put uptake occurred under different conditions of concentration and pH in African violet petals (*Saint pauliaionantha*), at a relatively lower concentration gradient (0.5–1.1 micromoles, pH = 5.0–5.5) and to a high

concentration gradient (from 0.1 to 100 millimolar, pH = 8.0) (Kakkar et al. 1998); therefore, PAs show the ability as buffers or regulators (Pandey et al. 2000).

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## 19.4 Polyamines in Stress Responses

There is creditable evidence confirming the role of PAs in stress tolerance. Scavenging of excess reactive oxygen species (ROS) is a direct strategy by which plants adapt to adverse environments. However, altering metabolism and accumulating beneficial metabolites, including PAs, are another ways of plants' defense mechanism. Often changes in PA metabolism and expression levels of their pathway genes are positively linked with enhanced tolerance of abiotic stresses in plants (Shi and Chan 2014). For example, overexpression of *FcWRKY70* resulted in Put accumulation to provide drought tolerance in *Fortunella crassifolia* (Gong et al. 2015). Spd promotes biomass accumulation and upregulates proteins involved in cell rescue. Spd was involved in inducing antioxidant enzymes in tomato (*Lycopersicon esculentum*) seedlings subjected to high-temperature conditions (Sang et al. 2017). Exogenous Spm treatment induced defense mechanism and caused resistance against a root rot pathogen, *Phytophthora capsici*, in *Capsicum annuum* (Koc et al. 2017). Furthermore, Spd was found to be vital for adjustment of intracellular PA pathways and endogenous PA homeostasis, which enhanced salt tolerance in rice (Saha and Giri 2017).

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## 19.5 Mode of Action of Polyamines in Stress Responses

PAs are reported to protect plants against stress through mechanisms including the following:

1. Polycationic PAs are known to bind anionic molecules such as nucleic acids and proteins, thus stabilizing them. This property may be important in preventing stress-induced damage to these macromolecules. Spd and Spm have been shown to prevent radiation and oxidative stress-induced strand breaks in DNA.
2. PAs are involved in the regulation of membrane transport in plants. They have been shown to block two slow and fast vacuolar cation channels. The effect of PAs is direct, and the channels open when PAs are withdrawn. PAs also affect vacuolar and plasma membrane  $H^+$  and  $Ca^{2+}$  pumps and have been reported to bring about stomatal closure in response to drought stress by blocking the activity of a KAT1-like inward  $K^+$  channel in the guard cell membrane. This is an indirect effect caused by low-affinity PA binding to the channel protein.
3. PAs are known to modulate ROS homeostasis. They are known for inhibiting metals auto-oxidation, thereby reducing the required electron supply for ROS generation. PAs are known to induce antioxidant enzymes in stress situations, as evidenced by use of inhibitors of PA biosynthesis or transgenics overexpressing enzymes involved in PA biosynthesis. For example, application of D-arginine,

which is an inhibitor of PA biosynthesis enzymes, resulted in reduced levels of PAs and increased levels of ROS. In addition, *ADC* overexpression influences increased tolerance to drought stress, causing reduced ROS generation in transgenic plants. PAs, especially Spm have roles as signaling molecules which activate the antioxidant machinery. Generation of H<sub>2</sub>O<sub>2</sub> by PA catabolism, which may be promoted when PA levels are above a specific threshold, is known to play an essential role in signaling cascade regulation during abiotic and biotic stress conditions. Additionally, abiotic stresses often induce accumulation of abscisic acid (ABA) and nitric oxide (NO) and the interactions between them. PAs trigger protective responses, including regulation of the channels for ion homeostasis and stomatal responses to enhance and maintain water content, thereby inducing the antioxidant machinery to check excessive ROS generation, with compatible osmolytes synthesis and accumulation. All of these phenomena occur to cascade the abiotic stress tolerance of plants (Shi and Chan 2014).

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## 19.6 Polyamines and Abiotic Stress

Polyamines modulate the plant's response to much broader range of abiotic stresses than expected, viz., drought, salinity, heavy metal toxicity, oxidative stress, chilling injury, high temperature, osmotic stress, water logging, and flooding tolerance as proved either by exogenous application of polyamines or by development of transgenic plants overexpressing the genes involved in polyamine biosynthesis (Gill and Tuteja 2010a, b). Increased or decreased levels of PAs either act as a signal or as a messenger (to transmit the perceived signals from the sensors) to articulate the plants' behavioral response spatially and temporally either to avoid or overcome stress. Modulated endogenous polyamine (free or conjugated or bound) levels are known to be involved in formation of polyamine-RNA complexes, thereby generating structural changes in RNA at physiological concentrations of potassium and magnesium ions (Igarashi and Kashiwagi 2000). Covalent linkage of polyamines to various enzymes or proteins (post-translational modification) involved in physiological processes under normal or stressed conditions was catalyzed by transglutaminase (TGase; EC 2.3.2.13) class of enzymes (Beninati et al. 1985; Folk 1980). Of various abiotic environmental stimuli under which polyamines get modulated and thereby its cellular functions were mineral nutrient deficiency (Richards and Coleman 1952; Coleman and Hegarty 1957), metal toxicity (Choudhary et al. 2012a, b), salinity (Lefèvre et al. 2001; Hummel et al. 2004), high (Oshima 2007) and low temperature (Hummel et al. 2004), drought (Bhatnagar et al. 2008; Alcazar et al. 2010), hypoxia (Moschou et al. 2008), osmotic (Lefèvre et al. 2001), and oxidative factors (Moschou et al. 2008; Bouchereau et al. 1999). PAs, also change ion channels (Takahashi and Kakehi 2010), stimulate special kind of protein synthesis, stimulate assembly of 30S ribosomal subunits, and stimulate Ile-tRNA formation (Igarashi and Kashiwagi 2000). Also, modulated titers of polyamines in combination with epibrassinolides, active form of brassinosteroids, were reported to regulate abscisic acid (ABA) and indole-3-acetic acid (IAA)

pathways which in turn enhances tolerance to metal toxicity (Choudhary et al. 2012a, b). PAs in combination with brassinosteroids besides modulating ABA and IAA pathways with their cascading effects for heavy metal tolerance also modulate levels of antioxidants like glutathione, ascorbic acid, proline, glycine-betaine, and antioxidant enzymes like glutathione reductase, peroxidase, catalase, and superoxide dismutase to cause stress tolerance (Choudhary et al. 2012a, b). Enhanced levels of polyamines either through exogenous feeding (Gill and Tuteja 2010a, b) or through heterologous expression of polyamine biosynthetic genes in transgenic plants (Liu et al. 2007) were reported to increase abiotic stress tolerance. However, the use of constitutively expressed promoters like CaMV35S, ubiquitin, and actin with polyamine biosynthetic genes toward stress tolerance may produce modulated polyamine levels even under normal conditions resulting in deleterious effects which causes reduced plant yield special concern toward agricultural crops (Katiyar et al. 1999).

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## 19.7 Polyamines Involvement in Regulation of Plant Stress Tolerance and Adaptation

Abiotic and biotic stresses cause alterations in the normal physiological processes of all plant organisms, including the economically important crops. Plant damage and productivity decrease are most often due to naturally occurring unfavorable factors of the environment—natural stress factors. Plant organisms are also imposed to a large scale with new stressors related to human activity—anthropogenic stress factors. Independent of the type of stress, an increased production of reactive oxygen species (ROS) occurs in plants that alter their normal physiological functions, decrease the biosynthetic capacity of plant organisms, and cause damage that may lead to plant death (Mittler 2002; Fujita et al. 2006; Ahmad et al. 2008; Gill and Tuteja 2010a, b; Potters et al. 2010). In plants, ROS are generated mainly as by-products of various processes requiring high metabolic activity or elevated electron flow by electron transport chains. The major targets of deleterious ROS action are cellular macromolecules as phospholipids, proteins, and nucleic acids. Plant organisms possess a complex of antioxidant protective systems in order to cope with destructive effects of the unfavorable environmental conditions. Beside enzymatic antioxidants and nonenzymatic antioxidants, a significant number of studies provided evidence that PAs also contribute to plant stress tolerance as a part of defense mechanisms or adaptation programs that help plants to mitigate the negative stress consequences. Since all stresses limit plant growth and crop productivity, the efforts of many scientists are focused to minimize the negative stress effects. A promising strategy to enhance plant tolerance and adaptation to an unfavorable environment is the use of a transgenic and molecular genetic approach to increase the cellular PA concentrations (Table 19.1).

**Table 19.1** Enhanced stress tolerance in transgenic plants engineered to overproduce polyamines (PAs) Source: Todorova et al. (2014)

Gene	Gene source	Transgenic plant	Increased tolerance	PAs overproduction
ADC	Oat	Rice	Salt tolerance	Put
ADC	Oat	Eggplant	Multiple abiotic stress tolerance, fungus wilt tolerance	Put, Spd (particularly conjugated forms), and free Spm fraction
ADC	Datura	Rice	Drought tolerance	Put, facilitating synthesis of Spm and Spd
ADC1	<i>Arabidopsis</i>	<i>Arabidopsis</i>	Freezing tolerance	Put
ADC2	<i>Arabidopsis</i>	<i>Arabidopsis</i>	Drought and cold tolerance	Put
PtADC	<i>Trifoliate orange</i>	Tobacco tomato	Improvement in dehydration and drought tolerance	Put
PtADC	<i>Trifoliate orange</i>	<i>Arabidopsis</i> adc 1-1 mutant	High osmoticum, drought, low temperature tolerance	Put
ODC	Mouse	Tobacco	Salinity tolerance	Put
SAMDC	Tritordeum	Rice	Salt tolerance	Spm and Spd
SAMDC	Human	Tobacco	Salt, drought, fungal wilt tolerance	Spd, put especially conjugated fraction
SAMDC	Carnation	Tobacco	Salt, cold, acidic, ABA tolerance	Put, Spd, and Spm
ySAMDC	Yeast	Tomato	High temperature tolerance	Spm and Spd after exposure to high temperature
MdSAMDC2	Apple	Tobacco	Low temperature, salt, osmotic tolerance	Free put, Spd, and Spm
SAMDC1	<i>Arabidopsis</i>	<i>Arabidopsis</i>	Salt, dehydration, ozone tolerance	Spm
MdSAMDC1	Apple	European pear	Salt, osmotic tolerance	Spd
MdSAMDC1	Apple	European pear	Multiple abiotic stress (salt, osmotic, Cu) tolerance	Spd
MdSAMDC1	Apple	European pear	Al tolerance	Spd
MdSAMDC1	Apple	European pear	Heavy metal (Cd, Pb, Zn) tolerance	Spd
MdSAMDC1	Apple	Tomato	Salinity tolerance	Spd and Spm
SPDS	Figleaf gourd	<i>Arabidopsis</i>	Chilling, freezing, salinity, hyperosmosis, drought, paraquat tolerance	Conjugated Spd

(continued)

**Table 19.1** (continued)

Gene	Gene source	Transgenic plant	Increased tolerance	PAs overproduction
SPDS (FSPD1)	Figleaf gourd	Sweet potato	Chilling heat	Spd
SAMS (SsSAMS2)	<i>Suaeda salsa</i>	Tobacco	Salt tolerance	Free Spm, Spd, and put

## 19.8 Abiotic Stress

### 19.8.1 Mineral Nutrient Deficiency

Since macro- and micronutrients are of paramount importance for normal plant growth and development, insufficient mineral nutrition affects all physiological processes in plants. To cope with mineral deficiency, plants usually augment PAs in response to nutrient shortage of potassium, boron, calcium, magnesium, and phosphorus (Wimalasekera et al. 2011). Boron deficiency caused an accumulation of free and conjugated Put and Spd (Camacho-Cristobal et al. 2005). Polyamines are also incremented by lower P-supply in roots and shoots of *Plantago lanceolata* L. (Bratek and Lang 2003). Plants grown on soils with low Ca concentration due to acidic deposition are exposed to multiple stress factors, and under these conditions, plants produce stress-related N-rich metabolites like Put, arginine,  $\gamma$ -aminobutyric acid, and proline in a species-particular manner for protection from toxic ammonia (Minocha et al. 2010). Potassium deficiency leads to increased levels of free PAs in *Betula pendula* and *Betula pubescens* (Sarjala and Kaunisto 2002) and cause a time-dependent pattern of Put accumulation during in vitro development of *Gentiana triflora* (Takahashi et al. 2012). Increased free-Put in potassium-deficient plants indicates that PAs are involved in maintenance of cation–anion balance in plant cells and conferred plant adaptation to ionic stress (Bouchereau et al. 1999).

### 19.8.2 Drought, Salt, Cold, and Osmotic Stress

Drought and salinity are widespread environmental constraints, and both cause reduced water potential and may lead to oxidative stress in plants (Lei 2008; Wang et al. 2011). The negative effects of salinity are deleterious involving complex mechanism, and they often come along ion toxicity. Drought increases accumulation of  $\text{Na}^+$  in cells and also results in ion toxicity (Wang et al. 2003; Basu et al. 2010). An increase of PAs due to drought and osmotic stresses has been found in different plant tissues (Yang et al. 2007; Lei 2008; Sziderics et al. 2010; Ghosh et al. 2011). Moderate or severe drought increased lipid peroxidation but also proline and PAs in wheat seedlings (Todorova et al. 2008). Frequently, PA accumulation (primarily Spd and/or Spm) is associated with drought and salt tolerance, supporting the effect of



PAs in response to stress and tolerance mechanisms (Basu et al. 2010; Zhou and Yu 2010; Alet et al. 2011, 2012). Mutlu and Bozcuk (2007) noted that the concentrations of free, bound, and total Spm increased in roots of *Helianthus annuus* L. (salt-tolerant and salt-sensitive) treated with NaCl and assumed its role in diminishing the harsh consequences of salinity. Salt tolerance in sunflower plants was related to the excessive accumulation of PAs in roots which were grown under salt stress conditions. Accumulation of Spm and compatible solutes (sucrose, proline, mannitol, and raffinose) was shown to be main players in alleviating NaCl stress in *Populus tremula* (Jouve et al. 2004). The role of higher PAs in osmotic adjustment as compatible solutes has been suggested (Basu et al. 2010; Alcazar et al. 2011a; Hussain et al. 2011).

Recently, Gupta et al. (2012a) reported that NaCl and Spd spray in rice (salt-tolerant and salt-sensitive) led to phosphorylation of 42-kDa  $\text{Ca}^{+2}$ -independent SnRK2 in roots of rice because of Spd in response to NaCl. Gupta et al. (2012b) demonstrated the important role of Spd in regulation of salinity-mediated signaling in rice. The response of plant varieties differing in tolerance to drought or salinity includes Put accumulation for sensitive cultivars and increase of Spm and Spd for tolerant varieties of rice (Roy et al. 2005; Roychoudhury et al. 2008; Basu et al. 2010), wheat (Liu et al. 2004b), and barley (Liu et al. 2006). Zhou and Yu (2010) demonstrated that rise of free and conjugated Spd and Spm in vetiver grass leaves assist plants to cope with water deficit conditions. Conversely, withholding water for 1 week in the drought-sensitive crop pepper caused significant accumulation in Put and cadaverine in leaves of pepper (Sziderics et al. 2010). Legocka and Sobieszczuk-Nowicka (2012) documented reduced amounts of microsome- and thylakoid-associated PAs in *Zea mays* and *Phaseolus vulgaris* subjected to iso-osmotic concentrations of NaCl and sorbitol. The authors suggested that these cultivars are drought sensitive and noted that PAs associated with microsomes and thylakoids might be good markers of plant stress tolerance. In order to show the effect of ionic and osmotic components of salinity on free PAs, Lefèvre et al. (2001) studied short-term exposure of salt-resistant and salt-sensitive rice cultivars to PEG or iso-osmotic concentrations of NaCl and KCl. The authors demonstrated that both ion and PA concentrations were changed 3 h after ionic stresses and assumed that the ionic component may trigger short-term PA accumulation independent of the osmotic component. The authors reported that enhanced Put levels also depends on the plant tissue in rice treated with PEG, KCl, or NaCl; it was found in the roots of salt-resistant cultivar, whereas in shoots for salt-sensitive cultivar. Similarly, Hummel et al. (2004) showed that the major effect, even after long-term exposure of *Pringlea antiscorbutica* seedlings to salinity and osmotic stress, was the modification of PA distribution between roots and shoots. Higher PA content in roots was assumed to be a developmental response to stress, and their accumulation in roots facilitated reinitiation of root growth. In contrast, salinity resistance in rice exposed to short-term PEG, KCl, and NaCl was not related with Spm or Spd (Lefèvre et al. 2001). A relationship between stress-induced augmentation of Spd and Spm in transgenic plants and tolerance to drought stress was also demonstrated. In addition, Peremarti et al. (2009) generated transgenic rice plants overexpressing *DsSAMDC* in order to

increase only higher PAs but not Put concentrations and noted accumulation of Spm, which facilitates drought recovery.

Alcazar et al. (2011b) analyzed *Arabidopsis* transcriptional profiles of PA biosynthetic genes and their metabolic fluxes through progressive drought acclimation with assessing the amounts of PAs in resurrection plant *Craterostigma plantagineum* and in PA biosynthetic mutants of *Arabidopsis*. The Put to Spm conversion in *Arabidopsis* coupled with Spm to Put back conversion has been proposed to participate in the drought stress response and to increase an effective PA recycling-loop during acclimation to water shortage. Involvement of Spm and Spd in maintaining ionic homeostasis by regulation of the activity of plasma membrane H<sup>+</sup>-ATPase and vacuolar H<sup>+</sup>-ATPase activities has been proposed as a general mechanism for salt tolerance and confirmed in different model systems (Roy et al. 2005; Liu et al. 2006; Janicka-Russak et al. 2010; Orsini et al. 2011). Similarly, due to experiments with PEG-treated drought-tolerant and drought-sensitive wheat cultivars, Liu et al. (2004a) reported that PAs conjugated to tonoplast vesicles correlate with maintenance of tonoplast H<sup>+</sup>-ATPase and H<sup>+</sup>-PPase activities in roots with enhanced osmotic stress tolerance in the plant. The authors could not detect Spm occurrence in wheat roots but free fractions of Spm- and Spd-enhanced osmotic stress tolerance in seedlings of the same cultivars (Liu et al. 2004b).

Yamaguchi et al. (2006, 2007) showed that the double knockout mutant of *Arabidopsis* (*acl5/spms*), which is unable to produce Spm, is hypersensitive to salinity and drought, but the addition of Spm specifically rescues the hypersensitive phenotype of mutant plants. The *acl5/spms* plant was shown to be Ca<sup>2+</sup> deficient (Yamaguchi et al. 2006), and as a result of improper stomatal closure under drought conditions, it lost more water as compared to the control (Yamaguchi et al. 2007). The authors proposed a model for the defensive role of Spm in salinity and drought stress responses (Yamaguchi et al. 2006; Kusano et al. 2007).

The expression of certain genes is also induced by ABA treatment (Urano et al. 2003). The *ADC2*, *SPDS1*, and *SPMS* expression was examined in ABA-deficient (*aba2-3*) and ABA-insensitive (*abi1-1*) mutants exposed to water stress exhibiting decreased transcriptional induction in the stressed *aba2-3* and *abi1-1* mutants compared to the wild type, showing that ABA modulates polyamine metabolism at the transcription level by upregulating the expression of *ADC2*, *SPDS1*, and *SPMS* genes under water stress conditions (Alcázar et al. 2006a, b). In addition, Put accumulation, occurred as a result of drought, is also damaged in the *aba2-3* and *abi1-1* mutants in comparison with wild-type plants. All these observations show that upregulation of PA-biosynthetic genes and accumulation of Put under water stress are especially due to ABA-dependent responses. It is assumed that polyamine responses to salt stress are also ABA-dependent, since both *ADC2* and *SPMS* are produced by ABA. In fact, stress-responsive, drought-responsive (DRE), low temperature-responsive (LTR), and ABA-responsive elements (ABRE and/or ABRE related motifs) are there in the promoters of the polyamine biosynthetic genes (Alcázar et al. 2006). This strengthens the view that in response to drought and salt treatments, the expression of some of the genes in polyamine biosynthesis are controlled by ABA. Free Put levels are enhanced on cold treatment, and this

corresponds with the introduction of *ADC* genes. Reduced expression of *NCED3* and several ABA-regulated genes was detected in the *adc1* mutants at low temperature. Complementation analyses of *adc1* mutants with ABA and mutual complementation of *aba2-3* mutant with Put supported the conclusion that diamine controls the levels of ABA in response to cold by modulating ABA biosynthesis at the transcriptional level (Cuevas et al. 2008, 2009). Certainly, Put and ABA are integrated in a positive feedback loop in which ABA and Put work synergistically for biosynthesis as a result of abiotic stress conditions that culminates a unique mode of operation of polyamines as regulators of ABA biosynthesis.

Liu et al. (2004a, b, c) reported that polyethylene glycol (PEG 6000) treatment (inducing water stress) significantly increased the free Spd and free Spm levels in the leaves of *Triticum aestivum* drought-tolerant cv. Yumai No. 18, whereas Yangmai No. 9 cv. (drought-sensitive) showed a significant increase of free Put. They suggested that free Spd, free Spm, and PIS-bound Put facilitated the osmotic stress tolerance of wheat seedlings. In response to PEG 6000 (20% w/v, 48 h)-mediated water stress condition in rice seedlings, Basu et al. (2010) showed that the salt-tolerant rice variety Pokkali accumulated the highest levels of Spd, Spm, and total PAs, while the Put level was the highest in the salt-sensitive variety IR-29 and induced maximally in the aromatic rice variety Pusa Basmati. In this respect, the aromatic variety behaved more closely to the sensitive variety. It was found that acquired tolerance to low water potential in potato cells leads to changes in Put biosynthesis and conjugation, which may be involved in ensuring cell survival (Gill and Tuteja 2010a, b). In case of *Theobroma cacao*, the expression of *TcODC*, *TcADC*, and *TcSAMDC* was induced with the onset of drought and correlated with the changes in stomatal conductance, photosynthesis, PSII efficiency, and leaf water potential. Induction of *TcSAMDC* in the leaves was more closely correlated with the changes in water potential (Bae et al. 2008). Yang et al. (2007) suggested that under drought conditions, the drought-resistant rice cultivars have the ability to respond early to water stress through increases in PA levels. Drought-resistant cultivars have higher SAMDC and SPDS activities and accumulated higher free Spd and free Spm in the leaves than drought-susceptible ones under water stress. The drought resistance in rice is therefore not only connected with PA levels but also with the response time at which PAs are significantly elevated. Vetiver grass can cope well with a moderate water-deficit conditions through maintenance of the total contents of free, conjugated, and bound Spd and Spm in leaves (Zhou and Yu 2010). In tolerant grapevine cultivars only, the higher PA-biosynthetic rate in the cellular compartment eliminates detrimental effects exerted by PAO-derived H<sub>2</sub>O<sub>2</sub> in the apoplast. In the sensitive genotype, intracellular homeostasis of PAs is not restored, and their levels are insufficient to mitigate the intervening effects of H<sub>2</sub>O<sub>2</sub> (Toumi et al. 2010). Endogenous levels of individual as well as total PAs in the roots of 7-day-old chickpea seedlings subjected to -0.8 MPa water stress increased significantly (Nayyar et al. 2005).

Using *Arabidopsis* has opened new vistas in functional dissection of the polyamine metabolic pathway and its role in the control of abiotic stress responses (Ferrando et al. 2004; Alcázar et al. 2006b; Kusano et al. 2008; Takahashi and

Takehi 2010; Gill and Tuteja 2010a). Annotation of the *Arabidopsis* genome enabled complete compilation for the biosynthetic pathway of polyamine. Transcript profiling by using Q-RT-PCR has revealed that water stress induces the expression of *ADC2*, *SPDS1*, and *SPMS* genes (Alcázar et al. 2006a). The expression of some of these genes is also induced by ABA treatment (Perez-Amador et al. 2002; Urano et al. 2003). To get a further insight into ABA regulation of polyamine pathway, the expression of *SPMS*, *SPDS1*, and *ADC2* was investigated in the ABA-deficient (*aba2-3*) and ABA-insensitive (*abi1-1*) mutants subjected to water stress (Alcázar et al. 2006a). These three genes display reduced transcriptional induction in the stressed *aba2-3* and *abi1-1* mutants compared to the wild type, showing that ABA modulates polyamine metabolism at the transcription level by upregulating the expression of *ADC2*, *SPDS1*, and *SPMS* genes under water stress conditions (Alcázar et al. 2006a). In addition, Put accumulation in response to drought is also hampered in the *aba2-3* and *abi1-1* mutants compared to wild-type plants. This result is further supported by metabolomic studies showing that polyamine responses to dehydration are also impaired in *nced3* mutants (Urano et al. 2009). All these observations support the conclusion that upregulation of PA-biosynthetic genes and accumulation of Put under water stress are mainly ABA-dependent responses. Under salt stress conditions, there is a rapid increase in the expression of *ADC2* and *SPMS*, which is maintained during the 24-h treatment and results in increased Put and Spm levels (Urano et al. 2003). Spm-deficient mutants are salt sensitive, while the introduction of Spm suppresses the salt sensitivity, suggesting a protective role of Spm to high salinity (Yamaguchi et al. 2006).

It is evident that polyamine responses to salt stress are also ABA-dependent, since both *ADC2* and *SPMS* are induced by ABA (see above). In fact, stress-responsive, drought-responsive (DRE), low temperature-responsive (LTR), and ABA-responsive elements (ABRE and/or ABRE-related motifs) are available in the promoters of the polyamine biosynthetic genes (Alcázar et al. 2006b). This reinforces the view that in response to drought and salt treatments, the expression of some of the genes involved in polyamine biosynthesis are regulated by ABA. Transcript profiling has also revealed that cold enhances the expression of *ADC1*, *ADC2*, and *SAMDC2* genes (Urano et al. 2003; Cuevas et al. 2008, 2009). Free Put levels are enhanced on cold treatment, and this corresponds with the induction of *ADC* genes. Surprisingly, the levels of free Spd and Spm remain unchanged or even decrease in response to cold treatment. The absence of correlation between enhanced *SAMDC2* expression and the decrease of Spm levels could be a result of increased Spm catabolism (Cuevas et al. 2008). Since double mutants completely devoid of ADC activity are not viable in *Arabidopsis* (Urano et al. 2005), two independent mutant alleles for both *ADC1* and *ADC2* were used to study their response to freezing. As indicated in “polyamines and abiotic stress,” the *adc1* and *adc2* mutations caused higher sensitivity to freezing conditions, in both acclimated and non-acclimated plants, while addition of Put complemented this stress sensitivity (Cuevas et al. 2008, 2009). Reduced expression of *NCED3* and several ABA-regulated genes was found in the *adc1* mutants at low temperature. Complementation analyses of *adc1* mutants with ABA and reciprocal complementation of

*aba2-3* mutant with Put supported the conclusion that this diamine controls the levels of ABA in response to cold by modulating ABA biosynthesis at the transcriptional level (Cuevas et al. 2008, 2009). All these results suggest that Put and ABA are integrated in a positive feedback loop, in which ABA and Put reciprocally promote each other's biosynthesis in response to abiotic stress. This highlights a novel mode of action of polyamines as regulators of ABA biosynthesis.

In general, cold-tolerant varieties show higher endogenous PA levels in response to low temperature than non-tolerant ones. Nayyar (2005) found that PA levels were enhanced by six to nine times in chickpea (*Cicer arietinum* L.) subjected to chilling temperatures. In poplar seedlings grown at 4 °C, Put accumulation occurred during the beginning of the cold treatment, while Spd and Spm accumulated after 4 or 7 days, respectively (Renaut et al. 2005). More recent quantitative expression analyses indicate that transcription of both *ADC1* and *ADC2* genes is produced as early as 30 min after cold exposure, the mRNA transcript levels of *ADC1* being higher than that of *ADC2*. The complementation analysis of *adc* mutants with ABA and reciprocal complementation tests of the *aba 2-3* mutant with Put help the research finding of Put controls ABA levels in response to low temperature (Cuevas et al. 2008). However, data obtained by Kim et al. (2002) in tomato showed that ABA and Put affected cold-induced changes in cellular membranes of tomato leaves independently. The sequence analysis of the *ADC1* promoter revealed the presence of CRT/DRE, which could mediate the early and transient *ADC1* upregulation under cold stress (Alcázar et al. 2006b). Accumulation of Put was also observed in lucerne and wheat during cold hardening, in which a differential regulation of *ADC* activity could be detected between control and cold-treated plants. In a chilling-tolerant rice cultivar (Tainung 67), the *ADC* activity and Put levels increased during low temperature in both shoots and roots, while a chilling-sensitive cultivar (Taichung Native 1) showed a slight Put rise in shoots and a lowering in roots (Lee et al. 1997). Chilling induced the expression of all the three *ADC* genes (*MADC1*, *MADC2*, and *MADC3*) in *Brassica juncea*, whereas salt predominantly resulted in increased accumulation of *MADC3* transcript (Mo and Pua 2002). The low temperature also increased the Put level in cold-sensitive maize plants, especially when the stress occurred in light. Pillai and Akiyama (2004) suggested that the induction of the *OsSAMDC* gene in response to cold could be utilized as a molecular marker for the ability of rice seedlings to withstand low temperatures. In *Arabidopsis*, the expression of *SAMDC2* also increased after cold treatment. Cuevas et al. (2008) reported absence of correlation between enhanced *SAMDC2* expression and decrease of Spm levels in response to cold treatment, which may be a result of increased Spm catabolism. Shen et al. (2000) showed that chilling markedly increased Spd content concomitantly with a rise in SAMDC activity in the cold-tolerant cucumber cultivars (Jinchun No. 3) but not in sensitive ones (Suyo). This response is not mediated by ABA, since Spd pretreatment did not affect the content of ABA in cold-treated leaves of the cv. Suyo, and ABA content did not increase in the leaves of the cv. Jinchun No. 3. Imai et al. (2004) reported that a novel *SPDS* gene, *OsSPDS2* from rice, was involved in chilling response in roots. This gene is closely related to *AtSPDS3* or *At5g53120*, a putative *Arabidopsis* *SPDS* gene. Overexpression of *ZAT12*, a cold-inducible gene

encoding a C2H2 zinc finger transcription factor (that contributes to an increased freezing tolerance), dampens the expression of the *CuAO* gene *At4G12280*, which shows sequence similarities to diamine oxidases. ZAT12 signaling promotes tolerance by activating a ZAT12 regulon, which includes *ADC1* and *ADC2* upregulation, leading to a resultant accumulation of Put (Alcázar et al. 2006b). Another cold inducible zinc finger protein called SCOF-1 has been reported from soybean (Kim et al. 2001).

Osmotic treatments using sorbitol induced high levels of Put and ADC in detached oat leaves (Flores and Galston 1984), whereas Spd and Spm show a dramatic decrease. Bouchereau et al. (1999) reported that osmotica with widely different assimilation routes, such as sorbitol, mannitol, proline, betaine, and sucrose, all induce a rise in Put. Such alterations are coincident with indications of stress, such as wilting and protein loss. Tiburcio et al. (1995) reported that when peeled oat leaves are incubated with sorbitol in the dark, they lose chlorophyll and senescence rapidly. Senescence could be slowed down by incorporating Spm in the incubation medium. The senescence-retarding effect of Spm was correlated with increase in the introduction of identified precursors into proteins, RNA, and DNA. They also inferred with rise in putrescine level and in particular its bound form to thylakoid membranes. A study by Schraudner et al. (1990) also discovered a correlation between ethylene emission, and PA biosynthesis was found in  $O_3^-$  treated potato and tobacco plants, of which the leaves showed early senescence in response to the pollutant. In the presence of  $O_3$ , all compounds of ethylene biosynthetic pathway in tobacco leaves were upregulated. Put and Spd levels also increased, as did ornithine decarboxylase (ODC) activity (Bouchereau et al. 1999). Differences in PA (Put, Spd, Spm) response under salt stress have been reported among and within species. For example, according to Prakash and Prathapsenan (1988), endogenous levels of PAs (Put, Spd, and Spm) decreased in rice seedlings under NaCl stress, whereas Basu et al. (1988) found that salinity influences accumulation of such compounds in the same material (Bouchereau et al. 1999).

Santa-Cruz et al. (1997) reported that the (Spd + Spm):Put ratios increased with salinity in the salt-tolerant tomato species (*Lycopersicon pennellii*, Carrel D'Arcy) but not in the salt-sensitive tomato species (*L. esculentum*). In both species, stress treatments decreased the levels of Put and Spd. The Spm levels did not decrease with salinity in *L. pennellii* over the salinization period; however, it was found to decrease in *L. esculentum*. The effects of different NaCl concentrations on maize embryogenic cells obtained from immature embryo cultures of a salt-sensitive inbred line (cv. w64) and a resistant hybrid (cv. Arizona) have also been reported where increased salt concentration strongly reduced the growth of the calluses and resulted in a significant increase in the total PA (Put, Spd) content, especially caused by a rise in Put. A study conducted by Bouchereau et al. (1999) revealed that incorporating the inhibitors of Put synthesis, the ADC pathway in tomato plants functions appropriately in both stress and control conditions, whereas the ODC pathway is stimulated only under the stress conditions.

## 19.9 Heavy Metals

Metal contamination of soils has considerably increased due to human activities, and heavy metal excess also provoked changes in PA metabolism. Abundance of PAs in different plants cultivated on abundant heavy metals and their effect in regulation of plant tolerance and adaptation to metal stress is well studied. Cadmium excess led to increase in Put, which was accompanied by a respective decrease in higher PAs in *Potamogeton crispus* L. (Yang et al. 2010). The activities of PAO and DAO increased categorically with the rise in Cd concentrations, and authors suggested that some PAs and their forms impart an important role in the adaptation mechanism of *P. crispus* under Cd-stress.

In experiments with tobacco BY-2 cells exposed to Cd, a marked accumulation of total PAs during 3-day treatment was observed (Kuthanová et al. 2004). The increase in PA levels was because of Put which was coincident with DAO activity. Cd treatment increased PAs in roots and leaves of *Salix alba* L., whereas in *Populus nigra* L., only the Put content was significantly enhanced in roots (Zacchini et al. 2011). Lefèvre et al. (2009) also observed that the halophyte *Atriplex halimus* exposed to Cd accumulated free PAs. Probably the resistance of *A. halimus* to Cd toxicity was related to the improved tissue tolerance through increase in the synthesis of osmoprotective compounds like PAs. Polyamine concentrations in mung bean plants grown on medium supplied with Cd was also higher than these found in control plants and negatively correlated with the activity of DAO (Choudhary and Singh 2000).

Increased contents of Ni, Zn, Cu, and Cd in soil increased the accumulation of Put and Spm in spinach and barley (Bergmann et al. 2001). Groppa et al. (2007a, b, 2008) found that PA metabolism was differently affected in wheat and sunflower discs subjected to either Cu or Cd. Put content increased significantly in sunflower plants after Cd and Cu treatments, Spd was modified only by Cd, while Spm was affected by Cu or Cd and rose at 16 days after the germination of seeds (Groppa et al. 2007b). Both ADC and ODC activities were increased by Cd, whereas Cu enhanced ADC but reduced ODC activity. Furthermore, Groppa et al. (2007a) studied the PA metabolism in wheat leaves under Cd or Cu stress and showed that cadmium and copper increased Put, Spd remained unaltered either with Cd or Cu, while Spm was reduced as compared to control values by both heavy metal treatments. Copper also decreased ADC but did not modify ODC and DAO activities (Groppa et al. 2007a). Both metals caused significant increases in proline and Put and also in Spd and Spm at the highest concentration tested (1 mM), and these increments were more pronounced for Cd treatment of sunflower seedlings (Groppa et al. 2008). Additionally, the authors suggested that PAs are key biological compounds, probably involved in signaling pathways triggered under stress environmental conditions. Similarly, Zhang et al. (2010) found that Cu treatment increased Put and lowered the Spd and Spm levels, thereby reducing the Put/(Spd + Spm) ratio in leaves of grafted and non-grafted cucumber seedlings. Grafting markedly reversed these Cu-induced effects for all Spd, Spr, and Put and partly regained the Put/(Spd + Spm) ratio in leaves. These results suggest that grafting can enhance the tolerance of cucumber

seedlings to Cu stress by increasing the activities of antioxidants and the levels of Spd and Spm, decreasing the Put/(Spd + Spm) ratio and the levels of ROS, promoting free proline and soluble protein synthesis in cucumber seedling leaves (Zhang et al. 2010).

In *Erica andevalensis* hydroponically grown with supplied increasing Cu concentration (1 to 500  $\mu\text{M}$ ) in the nutrient medium, the heavy metal led to a significant rise of PAs in shoots (Rossini Oliva et al. 2010). High levels of PAs in shoots helped the authors to conclude that PAs impart a defensive mechanism in plant cells at high metal concentration. Accordingly, Choudhary et al. (2009) showed that Cu treatment resulted in to a marked increase in PAs in *R. sativus* seedlings. Szafranska et al. (2011) examined the effect of Cu on the regeneration of *Daucus carota* L. androgenic embryos of var. Feria and 1014 breeding line and also on PAs, proline contents, lipid peroxidation, and Cu accumulation after 16 and 24 weeks and found that the higher tolerance of Feria to oxidative stress may result from increased content of proline and higher contents of the Put and Spd. The authors concluded that variations in PA levels depend not only on the concentrations of heavy metal but also on plant species and cultivars, and PAs point to better protection of such cultivars that possess higher constitutive PA levels. Zhao et al. (2008) investigated the phytotoxic effect of Ni on *Hydrocharis dubia* leaves and showed that Ni induced lipid peroxidation, destroyed the structure and functions of membranes, altered the balance of nutrient elements, and caused the toxicity of *H. dubia*. Ni stress significantly increased Put and lowered Spd and Spm levels, thus significantly reducing the ratio of free (Spd + Spm)/Put in leaves, which has been considered as a signal under stress. The changes of PAs content and physiological and biochemical responses in *H. dubia* leaves at high metal concentrations led the authors to conclude that PAs may be involved in the adaptation of plants to Ni-induced stress. Pb treatment of *Potamogeton crispus* L. significantly increased the free Put and changed dynamically other PAs levels.

The activities of PAO and ADC were initially decreased and then enhanced with the increase in the Pb concentration (Xu et al. 2011). Accumulation of PAs in shoot tissue of *Trifolium pratense* L. was affected by arsenic (Mascher et al. 2002) when metal was supplied in concentration 10 mg/kg soil. Extra accumulation of Spm seemed to be a part of a protective mechanism against the destruction of membranes caused by arsenate. In vitro shoots of a transgenic European pear, overexpressing *MaSPDS1*, performed attenuated susceptibility to heavy metal (Zn, Pb, Cu, Al, Cd) stress in conjunction with the wild plant (Wen et al. 2008, 2009, 2010). The authors supposed that abundance of Spd in transgenic plants could play an important role in the tolerance to heavy metals in long-term experiments. In this line Spd reduced the toxic effects of supplied metals mainly via alleviating oxidative status, by affecting mineral element balance and/or by its specific feature to act as metal chelator as well (Løvaas 1997; Lomozik et al. 2005). Thus, there is an indication that PAs acts as antioxidants and/or metal chelators (Kuthanová et al. 2004).



## 19.10 Herbicides

The extensive use of herbicides is a common practice in modern agriculture. However, their application usually provokes oxidative stress not only in sensitive but also in tolerant plants. Most of the herbicides are known to generate ROS by direct involvement in radical production or by inhibition of key biosynthetic pathways (Prasad and Rengel 1998). Enhancing free higher PAs in lotus prevented paraquat-induced superoxide production in vivo (Cuevas et al. 2004). However, because of the natural tolerance of *L. glaber* to the herbicide, it was not possible to correlate the content of free PAs with their hypothetical inhibitory role during oxidative stress. So, the authors tested for paraquat toxicity in tobacco plant and found that high concentrations of herbicide induced an increase of free PA levels in crude extracts and intercellular fluids, while low herbicide concentrations increased only the free PAs in the intercellular fluids. These results suggested that PA metabolism in the apoplast is involved in the physiological response to oxidative stress and subsequently in plant stress tolerance (Cuevas et al. 2004).

Deng (2005) found that chlorsulfuron and glyphosate directly regulated the levels of arginine and ornithine in *Datura stramonium* L. seedlings, which were increased two to three times. Recently, Kielak et al. (2011) investigated the effect of glyphosate-based herbicide (Roundup Ultra 360 SL) and showed significant positive correlation between Roundup dose and over-accumulation of Put and Spd as well as total PA in model plant duckweed. Putrescine was synthesized first within the plant cells and Spd later. Additionally, with augmentation of PAs, an increase in APX and CAT activities was recorded by herbicide which is the resultant of PA inclusion. These findings indicated that duckweed plants responded to herbicide stress by activating one or more components of the plant defense system, including PAs, which helps plants to mitigate the negative stress outcome.

## 19.11 Heat Tolerance

The amylose levels in rice are greatly decreased during elevated temperature conditions at the time of seed germination, changing the fine amylopectin structure and producing more chalky grains (Asaoka et al. 2010; Inouchi et al. 2000) which hampers the rice yield and seed quality (Krishnan and Ramakrishnan 2011; Peter 2011). The occurrence of chalky grains of rice is typically caused by the unusual expressions of genes encoding starch synthase enzymes (Nishi et al. 2001; Tanaka et al. 2010). As PAs are known to be involved in grain filling and they might contribute to build up heat resistance of some cereals, the hybrid rice “YLY 689” was taken in an experiment to find the effects of exogenous spermidine (Spd) on seed quality under high temperature during the grain filling stage (Fu et al. 2019). The concentration of Spm and Spd in superior grains of rice was remarkably higher than that of the inferior grains (Cao et al. 2016). In this investigation, just after pollination, spikes were treated with Spd or cyclohexylamine (CHA). Interestingly, CHA is its synthesis inhibitor.

Recently, Fu et al. (2019) revealed that when the rice plants were transferred to 40 °C for 5-day heat treatment, it resulted into significant enhancement in the germination index, percentage, germination index, seedling shoot height, vigor index, and dry weight of seeds in case of Spd pretreatment, harvested at 35 days after pollination. In contrast, the CHA significantly decreased the seed germination and seedling growth. Also, Spd categorically enhanced the peroxidase (POD) activity and reduced the malondialdehyde (MDA) content in seeds. Nevertheless, after spraying with Spd, the endogenous content of spermidine and spermine and the expression of their synthetic genes, *spermidine synthase (SPDSYN)*, and *spermine synthase (SPMS1 and SPMS2)* remarkably enhanced, although, the accumulation of amylose, total starch, and the expression of their related synthase genes, *soluble starch synthase II-3 (SS II-3)* and *granules bound starch synthase I (GBSSI)*, also rose to some magnitude indicating that exogenous Spd pretreatment could diminish the high temperature stress on rice seed grain filling and improve the rice seed quality, which could possibly be induced by upregulating endogenous polyamines and starch metabolism (Fu et al. 2019). Recent report also suggests that Spd is involved in heat tolerance in higher plants, which provides membrane stability and increase the reactive oxygen species (ROS) scavenging system. Cao et al. (2016) found that the content of Spd and Spm in superior grains of rice was significantly higher than that in inferior grains (Cao et al. 2016). The content of Spm and Spd in grains works in synergy with grain plumpness during the grain filling stage.

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## 19.12 Plant Senescence

The PA metabolic enzymes activities and PAs contents change throughout the stages of plant growth. In whole plants, endogenous PAs and PA synthetase activity were recorded to be highest in the meristem and growing cells and lowest in senescent tissues. As leaves senesce, the chlorophyll content slowly reduces and so does ADC and ODC activities, while PAO and hydrolases activities increase rapidly. All of these changes can be blocked by the inclusion of exogenous PAs (Duan 2000; Cai 2009). A reduction in PA levels seems to be a significant prelude to senescence signals, or it may be that a decrease in PAs content is the senescence signal (Duan et al. 2006). Exogenous Spd and Spm treatments can increase the content of PAs in cut flowers and prolong their senescence and improve quality (Yang and He 2001; Cao 2010). In *Anthurium andraeanum*, the introduction of GA<sub>3</sub> + Spm by spraying slowed down the senescence of cut flowers stored at 20 °C and augmented the quality of the inflorescences (Simões et al. 2018). Delayed leaf senescence was recorded to be linked with a rise in Spm concentration, reduced ROS production, and increased NO levels (Sobieszczuk-Nowicka 2017). Polyamines appeared to slow down the process of senescence by suppressing the ethylene biosynthesis (Woo et al. 2013; Anwar et al. 2015). 0.1 mM Spd application on gerbera flowers or gerbera vase flower fortification with 10 mM Spd resulted in slowing down the senescence, while those sprayed with 1 mM Spd, 10 mM Spd, 0.1 mM Spm, 1 mM Spm, or mixed solution of 0.1 mM each of Put, Spd, and Spm showed accelerated senescence, with brown spots and yellowing of the petal rims starting from day 2 of

treatment (Bagni and Tassoni 2006). Legocka and Sobieszczuk-Nowicka (2012) found that chlorophyll rapidly degraded, and Put accumulation inhibited protein degradation and reduced chlorophyll losses (Serafini-Fracassini et al. 2010; Cai et al. 2015). In peony, a PA synthesis inhibitor (0.1 Mm) extended the lifespan and delayed the cut flowers senescence, while PAs lessened the flower lifespan and accelerated flower senescence (Han 2016).

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### 19.13 Conclusion and Future Prospects

Many scientists have reported different plants defence mechanisms exhibited during its growth progression. However, the practical mode of action and regulations of many aspects still remain uncovered. The biosynthesis of PAs is present in every cell and tissue of plants; however, the number of enzymes involved in these processes is limited. That is why PA biosynthesis proves to be is an experimental model for studying how they act in protecting plants against various biotic and abiotic stresses. From several decades, PAs are being employed using mutants, inhibitors, plant tissue culture, exogenous application, genetic transformation techniques, etc. Various studies have been reported in order to illustrate the role of PAs such as Spm and Spd which have increased levels when plants get exposed to stress conditions enzymes, indicating the homeostatic equilibrium that is maintained inside the plant cells incorporating enzymes.

PAs' role in combating abiotic stress could easily be inferred by studies performed on *Arabidopsis*, wherein PAs were found to act antagonistically against salinity and cold stress conditions, thereby providing tolerance against such factors (Roy et al. 2005; Liu et al. 2006; Janicka-Russak et al. 2010; Orsini et al. 2011). Similarly, exogenous spray of Spd and Spm helped in delayed plant senescence in cut flowers and vase flowers (Yang and He 2001; Cao 2010). PA levels were found to be enhanced in plants experiencing mineral deficiencies such as potassium, boron, calcium, magnesium, and phosphorus (Wimalasekera et al. 2011). Likewise, there are reports from various researchers of how PAs play a pivotal role in plant stress conditions.

However, a lot had been studied and reported about PAs, but there still exists a lacuna of understanding the enzymes involved during biosynthesis of PAs, localization of PAs at cellular and subcellular levels, translocation and movements of free PAs, and their interaction with other plant hormones. Therefore, more research needs to be done under these niche areas so as to have a better understanding of polyamines as a whole.

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